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SEQUENCE LISTING

<110> INTERNATIONAL LIVESTOCK RESEARCH INSTITUTE THE INSTITUTE FOR GENOMIC RESEARCH

<120> EAST COAST FEVER VACCINE BASED ON CTL-SPECIFIC SCHIZONT ANTIGENS

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<150> 60/504,428

<151> 2003-09-22

<160> 53

<170> PatentIn Ver. 3.2

<210> 1

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Leu His Ser Pro Val Leu Gly Gly Asn Cys Ser His Glu Glu Leu Lys 20 25 30

Lys Leu Gly Met Leu Glu Gly Asp Gly Phe Asp Arg Asp Ala Leu Phe 35 40 45

Lys Ser Ser His Gly Met Gly Lys Val Gly Lys Arg Tyr Gly Leu Lys
50 60

Thr Thr Pro Lys Val Asp Lys Val Leu Ala Asp Leu Glu Thr Leu Phe 65 70 75 80

Gly Lys His Gly Leu Gly Gly Ile Ser Lys Asp Cys Leu Lys Cys Phe 85 90 95

Ala Gln Ser Leu Val Cys Val Leu Met Lys Cys Arg Gly Ala Cys Leu 100 105 110

Lys Gly Pro Cys Thr Asp Asp Cys Gln Asn Cys Phe Asp Arg Asn Cys
115 120 125

Lys Ser Ala Leu Leu Glu Cys Ile Gly Lys Thr Ser Ile Pro Asn Pro 130 135 140

Cys Lys Trp Lys Glu Asp Tyr Leu Lys Tyr Lys Phe Pro Glu Thr Asp 145 150 155 160

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Glu Asp Glu Ser Thr Lys Lys Gly Glu Ala Ser Gly Thr Ser 165 170

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Gln Phe Leu Lys Asn Val Ser Ala Leu Arg Arg Ser Ser Pro Asp Leu 20 25 30

Ser Pro Asp Gly Ser Phe Leu Gln Val Lys Ser Ala Ser Pro Gln Asp 35 40 45

Lys Gln Asp Val Ile Gln Ser Ser Pro Lys Val Thr Val Pro Thr
50 60

Val Asp Pro Glu Gly Leu Lys Lys Ala Val Thr Ala Ala Val Leu Ser
65 70 75 80

Asn Gln Asn Gln Ala Leu Gln Asn Gly Ala Leu Asn Pro Ala Asp Phe 85 90 95

Thr Gln Ala Ala Ser Val Asn Ser Met Ser Asn Ala Val Ser Ala Met
100 105 110

Asn Asn Thr Val Gly Pro Val Lys Asn Pro Met Ala Thr Val Gly Thr 115 120 125

Met Asn Ser Phe Thr Gly Met Pro Gly Val Gln Asp Asn Phe Pro Gln 130 135 140

Thr Pro Pro Val Asn Val Gln Asp Thr Ser Thr Gln Glu Asn Ser Leu 145 150 155 160

Asp Asn Leu Asn Leu Leu Leu Asp Pro Ser Leu Val Lys Ile Ser Gln 165 170 175

Ala Asp Ser His Ile Lys Glu Ser Met Glu Lys Ala Val His Ser Leu 180 185 190

Lys Lys Val Leu Glu Gly Leu Thr Asn Leu Ala Thr Leu Ser Lys Ser 195 200 205

Arg Asp Thr Glu Pro Phe Asn Val Leu Gly Asp Asp Tyr Thr Met Arg 210 215 220

Asn Val Leu Asp Leu Met Asn Lys Glu Leu Arg Gln Val Glu Ser Leu 225 230 235 240

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Gln Lys Val Val Phe Gln Phe Asn Ala Phe Ala Leu Ser Thr Phe Thr 245 250 255

Lys Ser Pro Asp Asp Asn Lys Lys Ser 260 265

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Ser Leu Tyr Asp Val Gly Ala Gly His Arg Ala Val Val Tyr Asn Arg
35 40 45

Ile Thr Gly Ile Ser Glu Thr Thr His Gly Glu Gly Thr His Phe Ile 50 55 60

Ile Pro Trp Leu Glu Arg Pro Ile Ile Tyr Asp Val Arg Thr Arg Pro 65 70 75 80

Arg Thr Leu Met Ser Leu Thr Gly Ser Arg Asp Leu Gln Met Val Asn 90 95

Ile Thr Cys Arg Val Leu Ser Arg Pro Asp Glu Arg Arg Leu Arg Asp
100 105 110

Ile Tyr Arg His Leu Gly Lys Asp Tyr Asp Glu Arg Val Leu Pro Ser 115 120 125

Ile Ile Asn Glu Val Leu Lys Ser Ile Val Ala Gln Tyr Asn Ala Ser 130 135 140

Gln Leu Ile Thr Gln Arg Glu Arg Val Ser Lys Ala Val Arg Asp Gln 145 150 155 160

Leu Val Asn Arg Ala Arg Asp Phe Asn Ile Leu Leu Asp Asp Val Ser 165 170 175

Leu Thr His Leu Ser Phe Ser Pro Glu Tyr Glu Lys Ala Val Glu Ala

Lys Gln Val Ala Gln Gln Gln Ala Glu Arg Ser Lys Tyr Ile Val Leu 195 200 205

Lys Ala Gln Glu Glu Lys Lys Ser Thr Ile Ile Lys Ala Gln Gly Glu 210 215 220

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Ser Glu Ala Ala Arg Leu Ile Gly Ser Ala Ile Lys Asp Asn Pro Ala
Phe Ile Thr Leu Arg Arg Ile Glu Thr Ala Lys Glu Val Ala Asn Ile
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Leu Ser Thr Asp Lys
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<210> 16
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representing amino acid residues #96 to amino acid #103

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ggtttcgaca gggatgcatt gttcaaatca tcacatggta tgggaaaggt aggaaaaagg 180
tatggtctta aaactactcc aaaagtagat aaagtcttag cagatcttga aacactgttt 240
ggaaaacacg gtcttggtgg tattagtaaa gattgtctta aatgttttgc acaaagccta 300
gtgtgcgtat taatgaaatg tagaggagca tgtctcaaag gaccatgtac tgacgactgc 360
caaaattgct ttgatagaaa ctgtaaatct gcattgctgg aatgcattgg gaaaacaagt 420
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gtaaaatcag cttctcctca ggataaacaa gatgtaatcc aaagttcctc tcctaaggta 180
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aatcccatgg ctactgttgg tactatgaac tcctttactg gaatgcctgg tgtacaggat 420
aattttcctc agacaccgcc tgttaatgtt caagacacct ctacccagga gaacagtctt 480
gacaacctaa atctcctctt agatccctcg ttagtaaaga tatctcaagc tgatagtcac 540
ataaaagaaa gcatggaaaa agctgtacac agccttaaaa aggtcttgga ggggctaacc 600
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tatacgatgc gtaacgtttt ggacctcatg aataaggaac tcaggcaggt tgaatctctt 720
cagaaagttg tgttccaatt caacgccttt gcactttcca ccttcactaa gagtccagac 780
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catagagctg ttgtatataa ccgtatcact ggaataagtg agactacaca tggagaagga 180
acgcacttca taattccctg gctagaacgt ccaataattt acgatgtgag gactcgtcct 240
aggactetga tgteteteae eggaageegt gaettgeaga tggttaacat cacetgeegt 300
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tacgacgage gagtectgee tteaataata aacgaggtte tgaagagtat tgtggcccag 420
tacaacgcct ctcagctcat tactcagaga gaaagagtta gcaaagcagt cagggaccag 480
ctggtgaaca gggccaggga ctttaatatt cttctcgatg atgtctcctt aacccactta 540
agetteagte etgaatatga aaaggetgta gaggetaaae aagtagetea acageaaget 600
gaacgcagta aatatatagt gttgaaggct caggaggaga agaaatcgac gataattaag 660
gctcagggag agtctgaggc tgcaaggctt attggaagtg caattaagga taaccctgcc 720
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